VENT& TOUCH LISTING

<110> Ryazanov, Alexey

<120> MAMMALIAN ALPHA-KINASE PROTEINS, NUCLEIC ACIDS AND DIAGNOSTIC AND THERAPEUTIC USES THEREOF

- <130> 601-1-098CIP
- <140> 09/832,292
- <141> 2001-04-10
- <150> 09/632,131
- <151> 2001-08-03
- <160> 45
- <170> PatentIn version 3.1
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- <211> 238
- <212> PRT
- <213> Homo sapiens
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Lys Arg Tyr Ile Glu Pro Val Asn Arg Asp Val Tyr Phe Glu Asp Val 50 55 60

Arg Leu Gln Met Glu Ala Lys Leu Trp Gly Asp Asp Tyr Asn Arg His 70 75 80

Lys Pro Pro Lys Gln Val Asp Ile Met Gln Met Cys Ile Ile Glu Leu 85 90 95

Lys Asp Arg Pro Gly Lys Pro Leu Phe His Leu Asp His Tyr Ile Asp 100 105 110

Gly Lys Tyr Ile Lys Tyr Asn Ser Asn Ser Gly Phe Val Arg Asp Asp 115 120 125

Asn Ile Arg Leu Thr Pro Gln Ala Phe Ser His Phe Thr Phe Glu Arg 130 135 140

Ser Gly His Gln Leu Ile Val Val Asp Ile Gln Gly Val Gly Asp Leu 145 150 155 160

Tyr Thr Asp Pro Gln Ile His Thr Glu Thr Gly Thr Asp Phe Gly Asn 165 170 175

Gly Asn Leu Gly Val Arg Gly Met Ala Leu Phe Phe Tyr Ser His Ala 180 185 190

Cys Asn Arg Ile Cys Glu Ser Met Gly Leu Ala Pro Phe Asp Leu Ser 195 200 205

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Tyr Ile Cys Gln Val Asp Arg Arg Val Leu Phe Asp Asp Val Arg Leu 50 55 60

Gln Met Asp Ala Lys Leu Trp Ala Glu Glu Tyr Asn Arg Tyr Asn Pro

70 75 80

Pro Lys Lys Ile Asp Ile Val Gln Met Cys Val Ile Glu Met Ile Asp 85 90 95

Val Lys Gly Ser Pro Leu Tyr His Leu Glu His Phe Ile Glu Gly Lys
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Tyr Ile Lys Tyr Asn Ser Asn Ser Gly Phe Val Ser Asn Ala Ala Arg 115 120 125

Leu Thr Pro Gln Ala Phe Ser His Phe Thr Phe Glu Arg Ser Gly His
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Gln Met Met Val Val Asp Ile Gln Gly Val Gly Asp Leu Tyr Thr Asp 145 150 155 160

Pro Gln Ile His Thr Val Val Gly Thr Asp Tyr Gly Asp Gly Asn Leu 165 170 175

Gly Ile Arg Gly Met Ala Leu Phe Phe His Ser His Arg Cys Asn Asp 180 185 190

Ile Cys Glu Thr Met Asp Leu Ser Asn Phe Glu Leu Ser Pro Pro Glu 195 200 205

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Phe Ala Glu Gly Ala Leu Arg Glu Ala Tyr His Thr Val Ser Leu Gly 20 25 30

Val Gly Thr Asp Glu Asn Tyr Asp Pro Leu Gly Thr Thr Thr Lys Leu 35 40 45

Phe Pro Pro Ile Glu Met Ile Ser Pro Ile Ser Lys Asn Asn Gly Ala 50 55 60

Met Thr Gln Leu Lys Asn Gly Thr Lys Phe Val Leu Lys Leu Tyr Lys 65 70 75 80

Lys Glu Ala Glu Gln Gln Ala Ser Arg Glu Leu Tyr Phe Glu Asp 'Val 85 90 95

Lys Met Gln Met Val Cys Arg Asp Trp Gly Asn Lys Phe Asn Gln Lys 100 105 110

Lys Pro Pro Lys Lys Ile Glu Phe Leu Met Ser Trp Val Val Glu Leu 115 120 125

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Ala Val Leu Thr Asn Arg Ser Thr Pro Gln Ala Phe Ser His Phe Thr 165 170 175

Tyr Gly Leu Ser Asn Lys Gln Met Ile Val Val Asp Ile Gln Gly Val 180 185 190

Asp Asp Leu Tyr Thr Asp Pro Gln Ile His Thr Pro Asp Gly Lys Gly 195 200 205

Phe Gly Leu Gly Asn Leu Gly Lys Ala Gly Ile Asn Lys Phe Ile Thr 210 225

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Lys Ser Gly Ala Ser Gly Arg Tyr Val Ser Lys Ile Gly Lys Lys Pro 35 40 45

Thr Pro Arg Pro Ser Tyr Phe Glu Asp Val Lys Met Gln Met Ile Ala 50 55 60

Lys Lys Trp Ala Asp Lys Tyr Asn Ser Phe Lys Pro Pro Lys Lys Ile 70 75 80

Glu Phe Leu Gln Ser Cys Val Leu Glu Phe Val Asp Arg Thr Ser Ser 90 95

Asp Leu Ile Cys Gly Ala Glu Pro Tyr Val Glu Gly Gln Tyr Arg Lys
100 105 110

Tyr Asn Asn Ser Gly Phe Val Ser Asn Asp Glu Arg Asn Thr Pro 115 120 125

Gln Ser Phe Ser His Phe Thr Tyr Glu His Ser Asn His Gln Leu Leu 130 135 140

Ile Ile Asp Ile Gln Gly Val Gly Asp His Tyr Thr Asp Pro Gln Ile 145 150 155 160

His Thr Tyr Asp Gly Val Gly Phe Gly Ile Gly Asn Leu Gly Gln Lys
165 170 175

Gly Phe Glu Lys Phe Leu Asp Thr His Lys Cys Asn Ala Ile Cys Gln 180 185 190

Tyr Leu Asn Leu Gln Ser Ile Asn Pro Lys Ser Glu Lys Ser Asp Cys 195 200 205

Gly Thr Val Pro 210

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<212> DNA

<213> Homo sapiens

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Asn Gln Asn Val Asn Ser Lys Val Asn Lys Tyr Tyr Ser Asn Leu Thr 50 55 60

Lys Ser Glu Arg Tyr Ser Ser Ser Gly Ser Pro Ala Asn Ser Phe His 70 75 80

Phe Lys Glu Ala Asn Lys His Ala Ile Gln Lys Ala Lys His Met Pro 85 90 95

Asp Pro Trp Ala Glu Phe His Leu Glu Asp Ile Ala Thr Glu Arg Ala 100 105 110

Thr Arg His Arg Tyr Asn Ala Val Thr Gly Glu Trp Leu Asp Asp Glu 115 120 125

Val Leu Ile Lys Met Ala Ser Gln Pro Phe Gly Arg Gly Ala Met Arg 130 135 140

Glu Cys Phe Arg Thr Lys Lys Leu Ser Asn Phe Leu His Ala Gln Gln 145 150 155 160

Trp Lys Gly Ala Ser Asn Tyr Val Ala Lys Arg Tyr Ile Glu Pro Val 165 170 175

Asp Arg Asp Val Tyr Phe Glu Asp Val Arg Leu Gln Met Glu Ala Lys 180 185 190

Leu Trp Gly Glu Glu Tyr Asn Arg His Lys Pro Pro Lys Gln Val Asp 195 200 205

Ile Met Gln Met Cys Ile Ile Glu Leu Lys Asp Arg Pro Gly Lys Pro 210 215 220

Leu Phe His Leu Glu His Tyr Ile Glu Gly Lys Tyr Ile Lys Tyr Asn 235 240

Ser Asn Ser Gly Phe Val Arg Asp Asp Asn Ile Arg Leu Thr Pro Gln 245 255

Ala Phe Ser His Phe Thr Phe Glu Arg Ser Gly His Gln Leu Ile Val

Val Asp Ile Gln Gly Val Gly Asp Leu Tyr Thr Asp Pro Gln Ile His 275 280 285

Thr Glu Thr Gly Thr Asp Phe Gly Asp Gly Asn Leu Gly Val Arg Gly 290 295 300

Met Ala Leu Phe Phe Tyr Ser His Ala Cys Asn Arg Ile Cys Glu Ser 305 310 315 320

Met Gly Leu Ala Pro Phe Asp Leu Ser Pro Arg Glu Arg Asp Ala Val 325 330 335

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Arg Pro Pro Leu Leu Arg Pro Leu Ser Glu Asn Ser Gly Asp Gly Asn 370 380

Met Ser Asp Val Thr Pro Asp Ser Leu Pro Ser Ser Pro Ser Ser Ala 385 390 395 400

Thr Pro His Ser Gln Lys Leu Asp His Leu His Trp Pro Val Phe Ser 405 410 415

Asp Leu Asp Asn Met Ala Ser Arg Asp His Asp His Leu Asp Asn His 420 425 430

Arg Glu Ser Glu Asn Ser Gly Asp Ser Gly Tyr Pro Ser Glu Lys Arg 435 440 445

Gly Glu Leu Asp Asp Pro Glu Pro Arg Glu His Gly His Ser Tyr Ser 450 455 460

Asn Arg Lys Tyr Gly Ser Asp Glu Asp Ser Leu Gly Ser Ser Gly Arg 465 470 475 , 480

Val Cys Val Glu Lys Trp Asn Leu Leu Asn Ser Ser Arg Leu His Leu 485 490 495

Pro Arg Ala Ser Ala Val Ala Leu Glu Val Gln Arg Leu Asn Ala Leu 500 505 510

Asp Leu Glu Lys Lys Ile Gly Lys Ser Ile Leu Gly Asp Val His Leu 515 520 525

Ala Met Val Arg Tyr His Glu Gly Gly Arg Phe Cys Glu Lys Gly Glu 530 540

Glu Trp Asp Gln Glu Ser Ala Val Phe His Leu Glu His Ala Ala Asn 545 550 550 560

Leu Gly Glu Leu Glu Ala Ile Val Gly Leu Gly Leu Met Tyr Ser Gln 565 570 575

Leu Pro His His Ile Leu Ala Asp Val Ser Leu Lys Glu Thr Glu Glu 580 585 590

Asn Lys Thr Lys Gly Phe Asp Tyr Leu Leu Lys Ala Ala Glu Ala Gly 595 600 605

Asp Arg Gln Ser Met Ile Leu Val Ala Arg Ala Phe Asp Ser Gly Gln 610 620

Asn Leu Ser Pro Asp Arg Cys Gln Asp Trp Leu Glu Ala Leu His Trp 625 630 635 640

Tyr Asn Thr Ala Leu Glu Met Thr Asp Cys Asp Glu Gly Gly Glu Tyr
645 650 655

Asp Gly Met Gln Asp Glu Arg Tyr Met Met Leu Ala Arg Glu Ala Glu 660 670

Met Leu Phe Thr Gly Gly Tyr Gly Leu Glu Lys Asp Pro Gln Arg Ser 675 680 685

Gly Asp Leu Tyr Thr Gln Ala Ala Glu Ala Ala Met Glu Ala Met Lys 690 695 700

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<212> DNA

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Asn Gln Asn Val Ser Ser Lys Val Gln Ser Tyr Tyr Ser Asn Leu Thr 50 55 60

Lys Thr Glu Leu Cys Gly Ser Thr Gly Ser Pro Ala Ser Ser Phe His 70 75 80

Phe Lys Glu Ala Trp Lys His Ala Ile Glu Lys Ala Lys His Met Pro 85 90 95

Asp Pro Trp Ala Glu Phe His Leu Glu Asp Ile Ala Thr Glu His Ala 100 105 110

Thr Arg His Arg Tyr Asn Ala Val Thr Gly Glu Trp Leu Lys Asp Glu 115 120 125

Val Leu Ile Lys Met Ala Ser Gln Pro Phe Gly Arg Gly Ala Met Arg 130 135 140

Glu Cys Phe Arg Thr Lys Lys Leu Ser Asn Phe Leu His Ala Gln Gln 145 150 155 160

Trp Lys Gly Ala Ser Asn Tyr Val Ala Lys Arg Tyr Ile Glu Pro Val 165 170 175

Asp Arg Ser Val Tyr Phe Glu Asp Val Gln Leu Gln Met Glu Ala Lys 180 185 190

Leu Trp Gly Glu Asp Tyr Asn Arg His Lys Pro Pro Lys Gln Val Asp 195 200 205

Ile Met Gln Met Cys Ile Ile Glu Leu Lys Asp Arg Pro Gly Gln Pro 210 220

Leu Phe His Leu Glu His Tyr Ile Glu Gly Lys Tyr Ile Lys Tyr Asn 235 240

Ser Asn Ser Gly Phe Val Arg Asp Asp Asn Ile Arg Leu Thr Pro Gln 245 250 255

Ala Phe Ser His Phe Thr Phe Glu Arg Ser Gly His Gln Leu Ile Val 260 265 270

Val Asp Ile Gln Gly Val Gly Asp Leu Tyr Thr Asp Pro Gln Ile His 275 280 285

Thr Glu Lys Gly Thr Asp Phe Gly Asp Gly Asn Leu Gly Val Arg Gly 290 295 300

Met Ala Leu Phe Phe Tyr Ser His Ala Cys Asn Arg Ile Cys Gln Ser 305 310 315 320

Met Gly Leu Thr Pro Phe Asp Leu Ser Pro Arg Glu Gln Asp Ala Val 325 330 335

Asn Gln Ser Thr Arg Leu Leu Gln Ser Ala Lys Thr Ile Leu Arg Gly 340 345 350

Thr Glu Glu Lys Cys Gly Ser Pro Arg Ile Arg Thr Leu Ser Ser Ser 355 360 365

Arg Pro Pro Leu Leu Leu Arg Leu Ser Glu Asn Ser Gly Asp Glu Asn 370 380

Met Ser Asp Val Thr Phe Asp Ser Leu Pro Ser Ser Pro Ser Ser Ala 385 390 395 400

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Val Cys Val Glu Thr Trp Asn Leu Leu Asn Pro Ser Arg Leu His Leu 485 490 495

Pro Arg Pro Ser Ala Val Ala Leu Glu Val Gln Arg Leu Asn Ala Leu 500 505 510

Asp Leu Gly Arg Lys Ile Gly Lys Ser Val Leu Gly Lys Val His Leu 515 520 525

Ala Met Val Arg Tyr His Glu Gly Gly Arg Phe Cys Glu Lys Asp Glu 530 540

Glu Trp Asp Arg Glu Ser Ala Ile Phe His Leu Glu His Ala Ala Asp 545 550 560

Leu Gly Glu Leu Glu Ala Ile Val Gly Leu Gly Leu Met Tyr Ser Gln 565 570 575

Leu Pro His His Ile Leu Ala Asp Val Ser Leu Lys Glu Thr Glu Glu 580 585 590

Asn Lys Thr Lys Gly Phe Asp Tyr Leu Leu Lys Ala Ala Glu Ala Gly 595 600 605

Asp Arg His Ser Met Ile Leu Val Ala Arg Ala Phe Asp Thr Gly Leu 610 620

Asn Leu Ser Pro Asp Arg Cys Gln Asp Trp Ser Glu Ala Leu His Trp 625 630 635 640

Tyr Asn Thr Ala Leu Glu Thr Thr Asp Cys Thr Glu Gly Gly Glu Tyr 645 650 655

Asp Gly Ile Gln Asp Glu Pro Gln Tyr Ala Leu Leu Ala Arg Glu Ala 660 665 670

Glu Met Leu Leu Thr Gly Gly Phe Gly Leu Asp Lys Asn Pro Gln Arg 675 680

Ser Gly Asp Leu Tyr Thr Gln Ala Ala Glu Ala Ala Met Glu Ala Met 690 695

Lys Gly Arg Leu Ala Asn Gln Tyr Tyr Gly Lys Ala Glu Glu Ala Trp 705 710

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Val Glu Lys Phe Asp His Gln Phe Asp Leu His Thr Gln Leu Met Ala 85 90 95

His Phe Thr Glu Gln Met Glu Asp Gln Leu Glu Lys Thr Met Lys Val

Val Arg Asn His Thr Asp Ser Leu Gly Gly Asn Val Gln Thr Lys Leu 115 120 125

Asp Glu Gly Ile Glu Lys Cys Met Ala Phe Ala Lys Lys Val Glu Gln 130 135 140

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Lys Asp Asn Arg Ser Glu Leu Glu Gly Leu Glu Lys Asp Cys Lys Asn 405 410 415

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Lys Leu Gln Asp Ser Ile Lys Arg Leu Glu Gln Asn Gln Lys Lys Ile 465 470 480

Glu Ala Glu Ile Gln Glu Gly Asn Glu Gln Val Glu Arg Val Leu Arg 485 490 495

Glu Glu Ala Ser Ile Ser Pro Ile Ser Ser Val Pro Lys Ser Pro Ile 500 505 510

Thr Thr Lys Arg Ser Ser Ile Ile Leu Asn Ser Pro Pro Met Thr Ser 515 520 525

Gln Gln Ser Ser Pro Lys Ile Gln Asp Leu Leu Ser Ser Ser Gly Ser 530 540

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Met Lys Leu Lys Val Glu Arg Lys Pro Phe Ala Glu Gly Ala Leu Arg 580 585 590

Glu Ala Tyr His Thr Val Ser Leu Gly Val Gly Thr Asp Glu Asn Tyr 595 600 605

Pro Leu Gly Thr Thr Lys Leu Phe Pro Pro Ile Glu Met Ile Ser 610 620

Pro Ile Ser Lys Asn Asn Glu Ala Met Thr Gln Leu Lys Asn Gly Thr 625 630 635 640

Lys Phe Val Leu Lys Leu Tyr Lys Lys Glu Ala Glu Gln Gln Ala Ser 650 655

Arg Glu Leu Tyr Phe Glu Asp Val Lys Met Gln Met Val Cys Arg Asp 660 670

Trp Gly Asn Lys Phe Asn Gln Lys Lys Pro Pro Lys Lys Ile Glu Phe 675 680 685

Leu Met Ser Trp Val Val Glu Leu Ile Asp Arg Ser Pro Ser Ser Asn 690 695 700

Gly Gln Pro Ile Leu Cys Ser Ile Glu Pro Leu Leu Val Gly Glu Phe 705 710 715 720

Lys Lys Asn Asn Ser Asn Tyr Gly Ala Val Leu Thr Asn Arg Ser Thr 725 730 735

Pro Gln Ala Phe Ser His Phe Thr Tyr Glu Leu Ser Asn Lys Gln Met 740 745 750

Ile Val Val Asp Ile Gln Gly Val Asp Asp Leu Tyr Thr Asp Pro Gln 755 760 765

Ile His Thr Pro Asp Gly Lys Gly Phe Gly Leu Gly Asn Leu Gly Lys 770 780

Ala Gly Ile Asn Lys Phe Ile Thr Thr His Lys Cys Asn Ala Val Cys 785 790 795 800

Ala Leu Leu Asp Leu Asp Val Lys Leu Gly Gly Val Leu Ser Gly Asn 805 810 815

Asn Lys Lys Gln Leu Gln Gln Gly Thr Met Val Met Pro Asp Ile Leu 820 825 830

Pro Glu Leu Met Pro Ser Asp Asn Thr Ile Lys Val Gly Ala Lys Gln 835 840 845

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- Lys Ser Ile Glu Ser Ile Ala Cys Asn Ser Asn Tyr Ile Phe Thr Ser 915 920 925
- Ser Pro Asp Asn Thr Ile Lys Val His Ile Ile Arg Ser Gly Asn Thr 930 935 940
- Lys Cys Ile Glu Thr Leu Val Gly His Thr Gly Glu Val Asn Cys Val 945 950 955 960
- Val Ala Asn Glu Lys Tyr Leu Phe Ser Cys Ser Tyr Asp Lys Thr Ile 965 970 975
- Lys Val Trp Asp Leu Ser Thr Phe Lys Glu Ile Lys Ser Phe Glu Gly 980 985 990
- Val His Thr Lys Tyr Ile Lys Thr Leu Ala Leu Ser Gly Arg Tyr Leu 995 1000 1005
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- Leu Ser Leu His Cys Thr Ala Ser Tyr Leu Phe Ser Thr Ser Lys 1040 1045
- Asp Asn Val Ile Lys Ile Trp Asp Leu Ser Asn Phe Ser Cys Ile 1055 1060 1065
- Asp Thr Leu Lys Gly His Trp Asn Ser Val Ser Ser Cys Val Val 1070 1075 1080
- Lys Asp Arg Tyr Leu Tyr Ser Gly Ser Glu Asp Asn Ser Ile Lys 1085 1090 1095
- Val Trp Asp Leu Asp Thr Leu Glu Cys Val Tyr Thr Ile Pro Lys 1100 1105 1110
- Ser His Ser Leu Gly Val Lys Cys Leu Met Val Phe Asn Asn Gln

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Leu Gln Ile Gly Lys Met Leu Pro Ser Gly Ser Gly Gly Gly Ala Thr 50 60

Ala Asp Ser Lys Phe Glu Lys Phe Lys Ala Arg Asn Thr Leu Ala Asp 65 70 75 80

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Pro Phe Ala Glu Gln Ala Phe Arg Lys Ala Tyr His Thr Leu Asp Leu 145 150 150

Ser Lys Ser Gly Ala Ser Gly Arg Tyr Val Ser Lys Ile Gly Lys Lys 165 170 175

Pro Thr Pro Arg Pro Ser Tyr Phe Glu Asp Val Lys Met Gln Met Ile 180 185 190

Ala Lys Lys Trp Ala Asp Lys Tyr Asn Ser Phe Lys Pro Pro Lys Lys 195 200 205

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Pro Gln Ser Phe Ser His Phe Thr Tyr Glu His Ser Asn His Gln Leu 260 270

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Ile His Thr Tyr Asp Gly Val Gly Phe Gly Ile Gly Asn Leu Gly Gln 290 295 300

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Gln Tyr Leu Asn Leu Gln Ser Ile Asn Pro Lys Ser Glu Lys Ser Asp 325 330 335

Cys Gly Thr Val Pro Arg Pro Asp Leu Ile Phe Pro Asp Thr Ser Glu

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Gln Tyr Leu Phe Ser Gly Ser Asn Asp Lys Ile Thr Lys Val Trp Asp 595 600 605

Leu Lys Thr Phe Arg Cys Asn Tyr Thr Leu Lys Gly His Thr Lys Trp 610 620

Val Thr Thr Ile Cys Ile Leu Gly Thr Asn Leu Tyr Ser Gly Ser Tyr 625 630 635 640

Asp Lys Thr Ile Arg Val Trp Asn Leu Lys Ser Leu Glu Cys Ser Ala 650 655

Thr Leu Arg Gly His Asp Arg Trp Val Glu His Met Val Ile Cys Asp 660 670

Lys Leu Leu Phe Thr Ala Ser Asp Asp Asn Thr Ile Lys Ile Trp Asp 675 680 685

Leu Glu Thr Leu Arg Cys Asn Thr Thr Leu Glu Gly His Asn Ala Thr 690 695 700

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<213> C. elegans

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Asn Asp Val Val Ile Glu Lys Pro Arg Met Asp Pro Leu His Val Arg 50 55 60

Lys Leu Met Glu Thr Trp Arg Lys Ala Ala Arg Arg Ala Arg Thr Asn 70 75 80

Tyr Ile Asp Pro Trp Lys Glu Phe Asn Ile His Glu Tyr Pro Val Gln 85 90 95

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Val Ala Met Glu Val Ala Ala Lys Gln Lys Lys Ser Cys Ile Val Pro 325 330 335

Pro Thr Val Phe Glu Ala Arg Arg Asn Arg Ile Ser Ser Glu Cys Val 340 345 350

His Val Glu His Gly Ile Ser Met Asp Gln Leu Arg Lys Arg Lys Thr 355 360 365

Leu Asn Gln Ser Ser Thr Asp Leu Ser Ala Lys Ser His Asn Glu Asp 370 380

Cys Val Cys Pro Glu Cys Ile Pro Val Val Glu Gln Leu Cys Glu Pro 385 390 395 400

Cys Ser Glu Asp Glu Glu Asp Glu Glu Glu Asp Tyr Pro Arg Ser Glu 405 410 415

Lys Ser Gly Asn Ser Gln Lys Ser Arg Arg Ser Arg Met Ser Ile Ser 420 430

Thr Arg Ser Ser Gly Asp Glu Ser Ala Ser Arg Pro Arg Lys Cys Gly 435 440 445

Phe Val Asp Leu Asn Ser Leu Arg Gln Arg His Asp Ser Phe Arg Ser 450 455 460

Ser Val Gly Thr Tyr Ser Met Asn Ser Ser Arg Gln Thr Arg Asp Thr 465 470 475 480

Glu Lys Asp Glu Phe Trp Lys Val Leu Arg Lys Gln Ser Val Pro Ala 485 490 495

Asn Ile Leu Ser Leu Gln Leu Gln Gln Met Ala Ala Asn Leu Glu Asn 500 505 510

Asp Glu Asp Val Pro Gln Val Thr Gly His Gln Phe Ser Val Leu Gly 515 520 525

Gln Ile His Ile Asp Leu Ser Arg Tyr His Glu Leu Gly Arg Phe Val 530 . 540

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Glu Lys His Thr Glu Gln Ser Pro Thr Asp Ala Tyr Gly Val Ile Asn 85 90 95

Phe Gln Gly Ser His Ser Tyr Arg Ala Lys Tyr Val Arg Leu Ser 100 105 110

Tyr Asp Thr Lys Pro Glu Val Ile Leu Gln Leu Leu Leu Lys Glu Trp 115 120 125

Gln Met Glu Leu Pro Lys Leu Val Ile Ser Val His Gly Gly Met Gln 130 135 140

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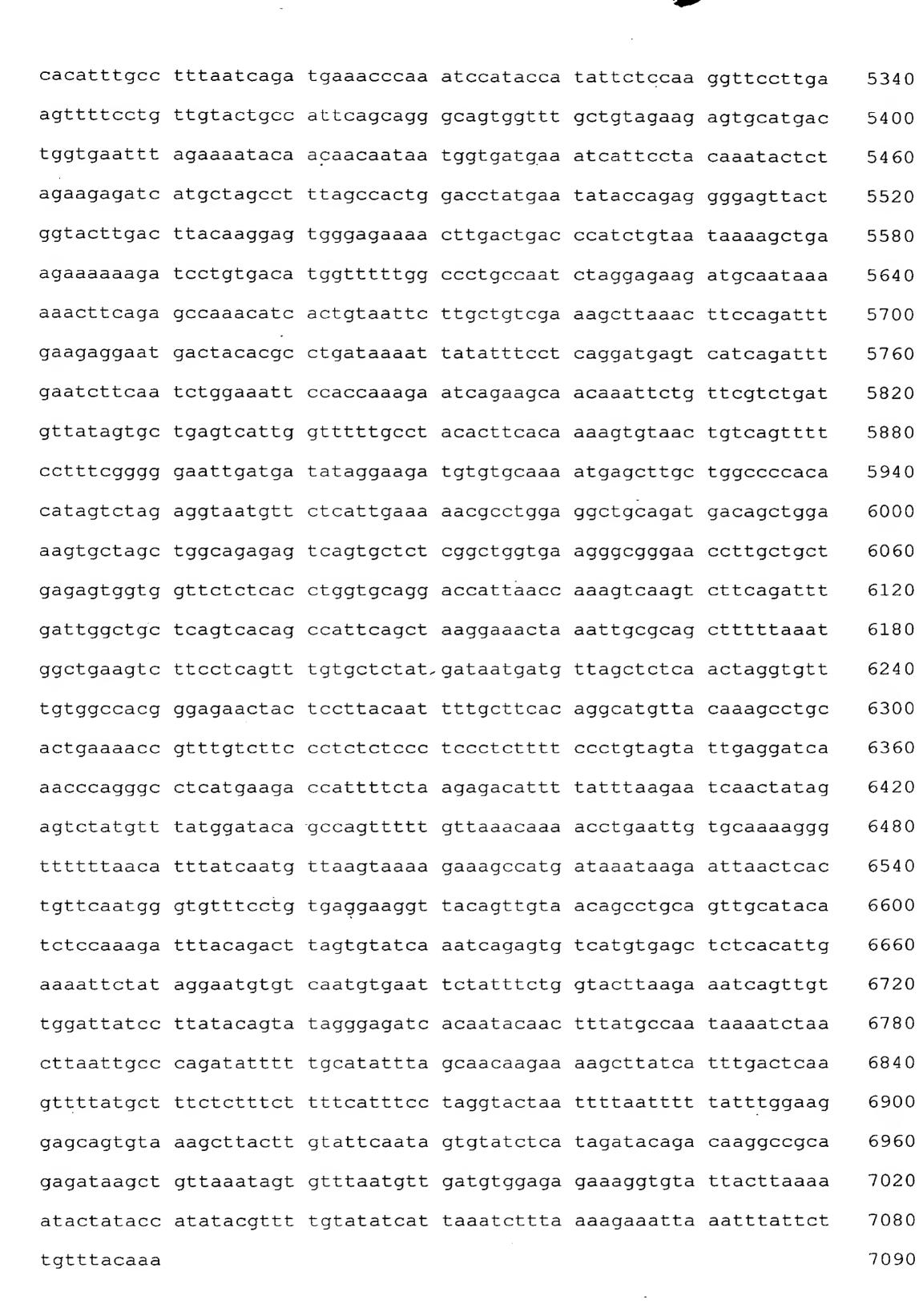
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Val Arg Leu Gly Glu His Phe Asn Gln Ala Ile Glu Glu Trp Ser Val 65 70 75 80

Glu Lys His Thr Glu Gln Ser Pro Thr Asp Ala Tyr Gly Val Ile Asn 85 90 95

Phe Gln Gly Ser His Ser Tyr Arg Ala Lys Tyr Val Arg Leu Ser 100 105 110

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Gln Met Glu Leu Pro Lys Leu Val Ile Ser Val His Gly Gly Met Gln 130 135 140

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Gln Arg Ile His Ala Arg Ile Gly Gln Gly Val Pro Val Val Ala Leu 275 280 285

Ile Phe Glu Gly Gly Pro Asn Val Ile Leu Thr Val Leu Glu Tyr Leu 290 295 300

Gln Glu Ser Pro Pro Val Pro Val Val Val Cys Glu Gly Thr Gly Arg 305 310 315 320

Ala Ala Asp Leu Leu Ala Tyr Ile His Lys Gln Thr Glu Glu Gly Gly 325 330 335

Asn Leu Pro Asp Ala Ala Glu Pro Asp Ile Ile Ser Thr Ile Lys Lys 340 345 350

Thr Phe Asn Phe Gly Gln Ser Glu Ala Val His Leu Phe Gln Thr Met 355 360 365

Met Glu Cys Met Lys Lys Glu Leu Ile Thr Val Phe His Ile Gly 370 375 380

Ser Glu Asp His Gln Asp Ile Asp Val Ala Ile Leu Thr Ala Leu Leu 385 390 395 400

Lys Gly Thr Asn Ala Ser Ala Phe Asp Gln Leu Ile Leu Thr Leu Ala 405 410 415

Trp Asp Arg Val Asp Ile Ala Lys Asn His Val Phe Val Tyr Gly Gln 420 425 430

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Met His Lys Phe Leu Thr Ile Pro Arg Leu Glu Glu Leu Tyr Asn Thr Lys Gln Gly Pro Thr Asn Pro Met Leu Phe His Leu Ile Arg Asp Val Lys Gln Gly Asn Leu Pro Pro Gly Tyr Lys Ile Thr Leu Ile Asp Ile Gly Leu Val Ile Glu Tyr Leu Met Gly Gly Thr Tyr Arg Cys Thr Tyr Thr Arg Lys Arg Phe Arg Leu Ile Tyr Asn Ser Leu Gly Gly Asn Asn Arg Arg Ser Gly Arg Asn Thr Ser Ser Ser Thr Pro Gln Leu Arg Lys Ser His Glu Thr Phe Gly Asn Arg Ala Asp Lys Lys Glu Lys Met Arg His Asn His Phe Ile Lys Thr Ala Gln Pro Tyr Arg Pro Lys Met Asp Ala Ser Met Glu Glu Gly Lys Lys Lys Arg Thr Lys Asp Glu Ile Val Asp Ile Asp Asp Pro Glu Thr Lys Arg Phe Pro Tyr Pro Leu Asn Glu Leu Leu Ile Trp Ala Cys Leu Met Lys Arg Gln Val Met Ala Arg Phe Leu Trp Gln His Gly Glu Glu Ser Met Ala Lys Ala Leu Val Ala Cys Lys Ile Tyr Arg Ser Met Ala Tyr Glu Ala Lys Gln Ser Asp Leu Val Asp Asp Thr Ser Glu Glu Leu Lys Gln Tyr Ser Asn Asp Phe Gly Gln Leu Ala Val Glu Leu Leu Glu Gln Ser Phe Arg Gln Asp Glu Thr Met

Ala Met Lys Leu Leu Thr Tyr Glu Leu Lys Asn Trp Ser Asn Ser Thr

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Gly Asp His Ala Gly Ile Asp Tyr Ser Trp Thr Ile Ser Ala Ala Lys 50 55 60

Gly Lys Glu Ser Glu Gln Trp Ser Val Glu Lys His Thr Thr Lys Ser 65 70 75 80

Pro Thr Asp Thr Phe Gly Thr Ile Asn Phe Gln Asp Gly Glu His Thr 85 90 95

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<212> PRT

<213> Homo sapiens

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Gly Ala Trp Ile Ile Thr Glu Gly Ile Asn Thr Val Ser Lys His Val
165 170 175

Gly Asp Ala Leu Lys Ser His Ser Ser His Ser Leu Arg Lys Ile Trp 180 185 190

Thr Val Gly Ile Pro Pro Trp Gly Val Ile Glu Asn Gln Arg Asp Leu 195 200 205

Ile Gly Lys Asp Val Val Cys Leu Tyr Gln Thr Leu Asp Asn Pro Leu 210 215 220

Ser Lys Leu Thr Thr Leu Asn Ser Met His Ser His Phe Ile Leu Ser 225 230 235 240

Asp Asp Gly Thr Val Gly Lys Tyr Gly Asn Glu Met Lys Leu Arg Arg 255

Asn Leu Glu Lys Tyr Leu Ser Leu Gln Lys Ile His Cys Arg Ser Arg 260 265 270

Gln Gly Val Pro Val Val Gly Leu Val Val Glu Gly Gly Pro Asn Val 275 280 285

Ile Leu Ser Val Trp Glu Thr Val Lys Asp Lys Asp Pro Val Val Val 290 295 300

Cys Glu Gly Thr Gly Arg Ala Ala Asp Leu Leu Ala Phe Thr His Lys 305 310 315 320

His Leu Ala Asp Glu Gly Met Leu Arg Pro Gln Val Lys Glu Glu Ile 325 330 335

Ile Cys Met Ile Gln Asn Thr Phe Asn Phe Ser Leu Lys Gln Ser Lys 340 345 350

His Leu Phe Gln Ile Leu Met Glu Cys Met Val His Arg Asp Cys Ile

Thr Ile Phe Asp Ala Asp Ser Glu Glu Gln Gln Asp Leu Asp Leu Ala Ile Leu Thr Ala Leu Leu Lys Gly Thr Asn Leu Ser Ala Ser Glu Gln Leu Asn Leu Ala Met Ala Trp Asp Arg Val Asp Ile Ala Lys Lys His Ile Leu Ile Tyr Glu Gln His Trp Lys Pro Asp Ala Leu Glu Gln Ala Met Ser Asp Ala Leu Val Met Asp Arg Val Asp Phe Val Lys Leu Leu Ile Glu Tyr Gly Val Asn Leu His Arg Phe Leu Thr Ile Pro Arg Leu Glu Glu Leu Tyr Asn Thr Lys Gln Gly Pro Thr Asn Thr Leu Leu His His Leu Val Gln Asp Val Lys Gln His Thr Leu Leu Ser Gly Tyr Arg Ile Thr. Leu Ile Asp Ile Gly Leu Val Val Glu Tyr Leu Ile Gly Arg Ala Tyr Arg Ser Asn Tyr Thr Arg Lys His Phe Arg Ala Leu Tyr Asn Asn Leu Tyr Arg Lys Tyr Lys His Gln Arg His Ser Ser Gly Asn Arg Asn Glu Ser Ala Glu Ser Thr Leu His Ser Gln Phe Ile Arg Thr Ala Gln Pro Tyr Lys Phe Lys Glu Lys Ser Ile Val Leu His Lys Ser Arg Lys Lys Ser Lys Glu Gln Asn Val Ser Asp Asp Pro Glu Ser Thr Gly Phe Leu Tyr Pro Tyr Asn Asp Leu Leu Val Trp Ala Val Leu Met Lys

Arg Gln Lys Met Ala Met Phe Phe Trp Gln His Gly Glu Glu Ala Thr 610 620

Val Lys Ala Val Ile Ala Cys Ile Leu Tyr Arg Ala Met Ala His Glu 625 630 635 640

Ala Lys Glu Ser His Met Val Asp Asp Ala Ser Glu Glu Leu Lys Asn 645 650 655

Tyr Ser Lys Gln Phe Gly Gln Leu Ala Leu Asp Leu Leu Glu Lys Ala 660 670

Phe Lys Gln Asn Glu Arg Met Ala Met Thr Leu Leu Thr Tyr Glu Leu 675 680 685

Arg Asn Trp Ser Asn Ser Thr Cys Leu Lys Leu Ala Val Ser Gly Gly 690 700

Leu Arg Pro Phe Val Ser His Thr Cys Thr Gln Met Leu Leu Thr Asp 705 710 715 720

Met Trp Met Gly Arg Leu Lys Met Arg Lys Asn Ser Trp Leu Lys Ile 725 730 735

Ile Ile Ser Ile Ile Leu Pro Pro Thr Ile Leu Thr Leu Glu Phe Lys 740 745 750

Ser Lys Ala Glu Met Ser His Val Pro Gln Ser Gln Asp Phe Gln Phe 755 760 765

Met Trp Tyr Tyr Ser Asp Gln Asn Ala Ser Ser Ser Lys Glu Ser Ala 770 780

Ser Val Lys Glu Tyr Asp Leu Glu Arg Gly His Asp Glu Lys Leu Asp 785 790 795 800

Glu Asn Gln His Phe Gly Leu Glu Ser Gly His Gln His Leu Pro Trp 805 810 815

Thr Arg Lys Val Tyr Glu Phe Tyr Ser Ala Pro Ile Val Lys Phe Trp 820 825 830

Phe Tyr Thr Met Ala Tyr Leu Ala Phe Leu Met Leu Phe Thr Tyr Thr 835 840 845

- Val Leu Val Glu Met Gln Pro Gln Pro Ser Val Gln Glu Trp Leu Val 850 855 860
- Ser Ile Tyr Ile Phe Thr Asn Ala Ile Glu Val Val Arg Glu Val Ser 865 870 875 880
- Ile Ser Glu Pro Gly Lys Phe Thr Gln Lys Val Lys Val Trp Ile Ser 890 895
- Glu Tyr Trp Asn Leu Thr Glu Thr Val Ala Ile Gly Leu Phe Ser Ala 900 905 910
- Gly Phe Val Leu Arg Trp Gly Asp Pro Pro Phe His Thr Ala Gly Arg 915 920 925
- Leu Ile Tyr Cys Ile Asp Ile Ile Phe Trp Phe Ser Arg Leu Leu Asp 930 935 940
- Phe Phe Ala Val Asn Gln His Ala Gly Pro Tyr Val Thr Met Ile Ala 945 950 955 960
- Lys Met Thr Ala Asn Met Phe Tyr Ile Val Ile Ile Met Ala Ile Val 965 970 975
- Leu Leu Ser Phe Gly Val Ala Arg Lys Ala Ile Leu Ser Pro Lys Glu 980 985 990
- Pro Pro Ser Trp Ser Leu Ala Arg Asp Ile Val Phe Glu Pro Tyr Trp 995 1000 1005
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- Leu Ile Ala Phe Phe Asn Asn Val Tyr Leu Asp Met Glu Ser Ile 1055 1060 1065
- Ser Asn Asn Leu Trp Lys Tyr Asn Arg Tyr Arg Tyr Ile Met Thr 1070 1075 1080
- Tyr His Glu Lys Pro Trp Leu Pro Pro Pro Leu Ile Leu Leu Ser

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Tyr						Ser 1270					Leu 1275	Ala	Gly	Gly
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Ser Gly Ser Ser Glu Ile Gly Gln Gly Ala Trp Val Lys Ala Lys 1550 1555 1560 Met Leu Thr Lys Asp Arg Arg Leu Ser Lys Lys Lys Asn Thr 1565 1570 1575 Gln Gly Leu Gln Val Pro Ile Ile Thr Val Asn Ala Cys Ser Gln 1580 1585 1590 Ser Asp Gln Leu Asn Pro Glu Pro Gly Glu Asn Ser Ile Ser Glu 1600 1595 1605 Glu Glu Tyr Ser Lys Asn Trp Phe Thr Val Ser Lys Phe Ser His 1615 1610 1620 Thr Gly Val Glu Pro Tyr Ile His Gln Lys Met Lys Thr Lys Glu 1625 1630 1635

Ile Gly Gln Cys Ala Ile Gln Ile Ser Asp Tyr Leu Lys Gln Ser 1640 1645 1650

Gln Glu Asp Leu Ser Lys Asn Ser Leu Trp Asn Ser Arg Ser Thr 1655 1660 1665

Asn Leu Asn Arg Asn Ser Leu Leu Lys Ser Ser Ile Gly Val Asp 1670 1680

Lys Ile Ser Ala Ser Leu Lys Ser Pro Gln Glu Pro His His His 1685 1690 1695

Tyr Ser Ala Ile Glu Arg Asn Asn Leu Met Arg Leu Ser Gln Thr 1700 1705 1710

Ile Pro Phe Thr Pro Val Gln Leu Phe Ala Gly Glu Glu Ile Thr 1715 1720 1725

Val Tyr Arg Leu Glu Glu Ser Ser Pro Leu Asn Leu Asp Lys Ser 1730 1740

Met Ser Ser Trp Ser Gln Arg Gly Arg Ala Ala Met Ile Gln Val 1745 1750 1755

Leu Ser Arg Glu Glu Met Asp Gly Gly Leu Arg Lys Ala Met Arg 1760 1765 1770

Val Val Ser Thr Trp Ser Glu Asp Asp Ile Leu Lys Pro Gly Gln

1775 1780 1785

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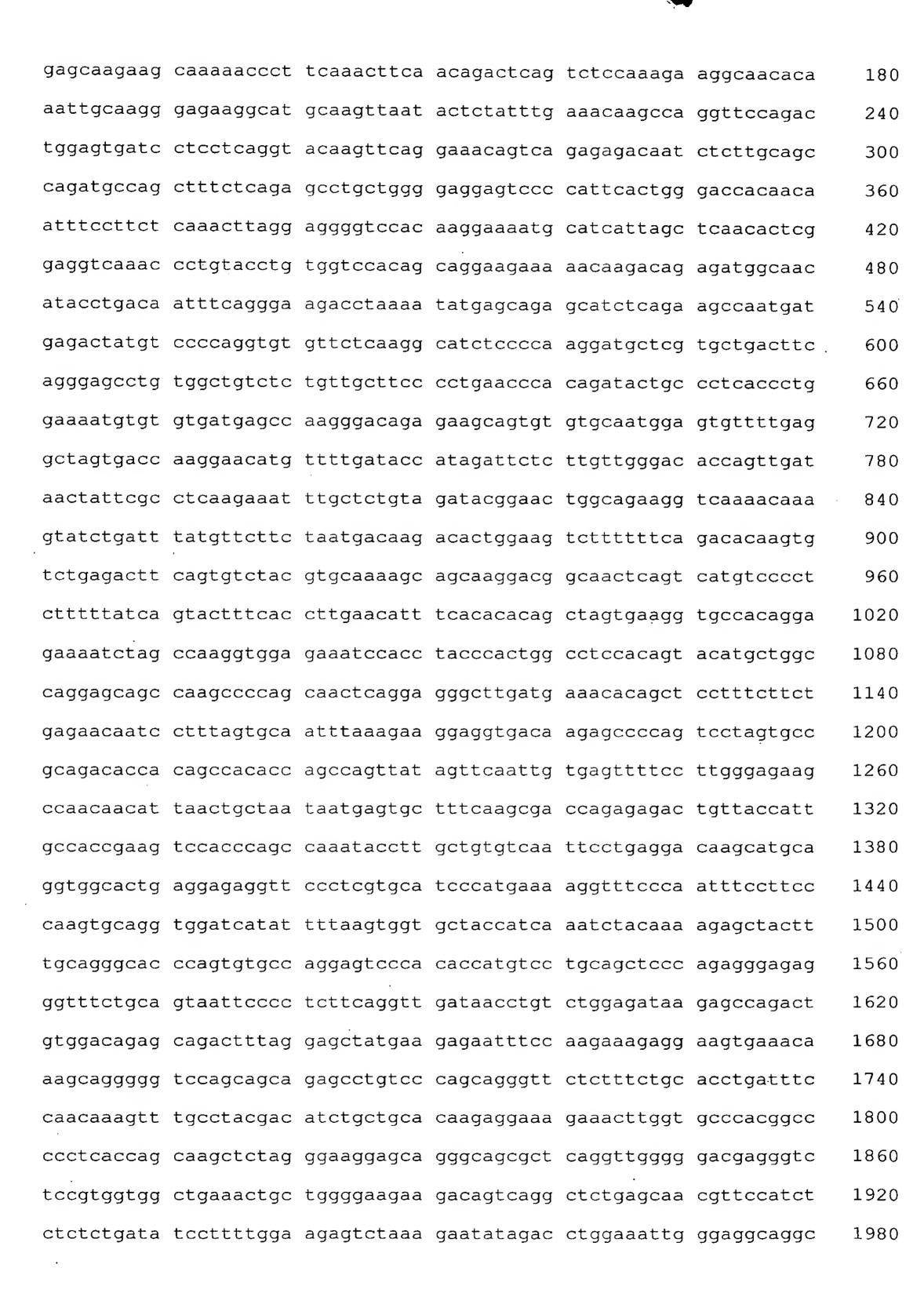
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Ile Ser Cys Ser Gln Met Pro Ala Phe Ser Glu Pro Ala Gly Glu Glu 100 105 110

Ser Pro Phe Thr Gly Thr Thr Thr Ile Ser Phe Ser Asn Leu Gly Gly 115 120 125

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Asp Lys Thr Leu Glu Val Phe Phe Gln Thr Gln Val Ser Glu Thr Ser 290 295 300

Val Ser Thr Cys Lys Ser Ser Lys Asp Gly Asn Ser Val Met Ser Pro 305 310 315 320

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Ala Thr Arg Glu Thr Val Thr Ile Ala Thr Glu Val His Pro Ala Lys 435 440 445

Tyr Leu Ala Val Ser Ile Pro Glu Asp Lys His Ala Gly Gly Thr Glu 450 455 460

Glu Arg Phe Pro Arg Ala Ser His Glu Lys Val Ser Gln Phe Pro Ser 465 470 475 480

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Lys Lys Phe Leu Thr Cys Pro Lys Ile Leu Glu Ser Ser Val Asp Pro 820 825 830

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Val Pro Cys Leu Gln Gly Thr Ile Leu Ser Glu Asn Arg Ile Ser Arg 885 890 895

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Gly Ala Asp Gln Val Ser Cys Glu Ala Arg Asp Glu Asp Asn Ser Gln 340 345

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Gly 385	Asp	His	Gly	His	Phe 390	Ser	Met	Pro	Glu	Gly 395	Gln	Gly	Leu	Cys	Ser 400

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Glu Gly Ala His Ser Arg Gly Leu Glu Glu His Phe Gln Glu Lys Gly 420 425 430

Ser Gly Met Lys His Gly Ile Arg Pro Gln Ser Thr Ser His Gln Val 435 440 445

Ser Leu Ser Ala Asn Asp Phe Gln Glu Ile Leu Pro Ser Ile Pro Thr 450 455 460

Met Gln Gln Glu Thr Asn Val Glu Pro Leu Glu His Ser Leu Ala Asp 465 470 475 480

Ser Arg Glu Glu Ile Glu Cys Ser Ser Asp Pro Arg Thr Ser Asp Leu 485 490 495

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Leu Gly Ser Trp Ala Val Gly Gly Lys Val Lys Ile Ile Thr Leu Glu 530 540

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Tyr Lys Glu Ala Glu Ala Gly Leu Thr Met Pro Gly Arg Ser Trp Ala 565 570 575

Leu Ser Asp Ile Leu Arg Ala Gly Ala Thr Arg Ser Glu Pro Gly Ala 580 585 590

Leu Gly Gly Ala Ala Trp Val Pro Ser Pro Gln Ala Asp Ala Leu Met 595 600 605 .

Ala Leu Gly Ala Asn Arg Asp Thr Trp Leu Gly Ala Ala Pro Asp Arg

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Gln Ala Asn Cys Asn Cys Leu Ser Ser Gln Cys Leu Ser Gln Pro Arg 625 630 635 640

Phe Leu Glu Ser Ser Val Asp Pro Val Glu Asp Lys Glu Leu Glu Val 655

Thr Asp Ser Pro Ser Glu Val Ser Lys Thr Gly Glu Met Glu Met Pro 660 670

Glu Thr Leu Asn Glu Glu Glu Glu Glu Thr Gln Gln Met Leu Arg His 675 680 685

Pro Ala Val Val Asn Gln Ser Val Asn Phe Pro Arg Ile Leu Glu Ser 690 700

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Glu Lys Pro Glu Pro Ser Asp Ser Ser Val Glu Gly Asn Glu Pro Ile 725 730 735

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Ser Cys Leu Thr Gly Val Thr His Thr Cys Val Lys Ala Glu Thr Asn 850 860

Asn Ser Thr Gly His Ile Tyr Gly Gly Ser Glu Pro Arg Thr Arg Gln 865 870 875 880

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Gly Asn His Val Arg Ser Ser Asp Asp Leu Thr Asn Thr Pro Cys Thr 900 910

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Gln Lys Pro Val Ala Ala Gln Val Ser Lys Ala Pro Gly Asp Gln Ser 995 1000 1005

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Gly Ser Gly His Leu Thr Ala Gly Ile Lys Lys Lys Ile Leu Ser 1025 1030 1035

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1325 1330 1335

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Asp Leu Ala Pro Leu Ala Pro Ala Arg Pro Arg Gly Pro Leu Ile Cys 50 55

His Thr Gly His Glu Gln Ala Gly Arg Glu Pro Gly Pro Gly Ser Ser 65 . 75 . 80

Thr Lys Gly Pro Val Leu His Asp Gln Asp Thr Arg Cys Ala Phe Leu 85 90 95

Pro Arg Pro Pro Gly Pro Leu Gln Thr Arg Arg Tyr Cys Arg His Gln
100 105 110

Gly Arg Gln Gly Ser Gly Leu Gly Ala Gly Pro Gly Ala Gly Thr Trp 115 120 125

Ala Pro Ala Pro Pro Gly Val Ser Lys Pro Arg Cys Pro Gly Arg Ala 130 135 140

Arg Pro Gly Glu Gly Gln Gln Gln Val Thr Thr Ala Arg Pro Pro Ala 145 150 155 160

Ile Asn Arg Gly Ala Arg Gln Pro Arg Ala Gly Ala Ala Ala Gly 165 170 175

Arg Gly Pro Gly Ala Gly Ala Trp Arg Thr Gly Glu Ala Ala Ala Ser 180 185 190

Ala Gly Pro Ala Val Gly Glu Gly Gly Ala Met Gly Ser Arg Arg Ala 195 200 205

Pro Thr Arg Gly Trp Gly Ala Gly Gly Arg Ser Gly Ala Gly Gly Asp 210 215 220

Gly Glu Asp Asp Gly Pro Val Trp Ile Pro Ser Pro Ala Ser Arg Ser Tyr Leu Leu Ser Val Arg Pro Glu Thr Ser Leu Ser Ser Asn Arg Leu Ser His Pro Ser Ser Gly Arg Ser Thr Phe Cys Ser Ile Ile Ala Gln Leu Thr Glu Glu Thr Gln Pro Leu Phe Glu Thr Thr Leu Lys Ser Arg Ser Val Ser Glu Asp Ser Asp Val Arg Phe Thr Cys Ile Val Thr Gly Tyr Pro Glu Pro Glu Val Thr Trp Tyr Lys Asp Asp Thr Glu Leu Asp Arg Tyr Cys Gly Leu Pro Lys Tyr Glu Ile Thr His Gln Gly Asn Arg His Thr Leu Gln Leu Tyr Arg Cys Arg Glu Glu Asp Ala Ala Ile Tyr Gln Ala Ser Ala Gln Asn Ser Lys Gly Ile Val Ser Cys Ser Gly Val Leu Glu Val Gly Thr Met Thr Glu Tyr Lys Ile His Gln Arg Trp Phe Ala Lys Leu Lys Arg Lys Ala Ala Ala Lys Leu Arg Glu Ile Glu Gln Ser Trp Lys His Glu Lys Ala Val Pro Gly Glu Val Asp Thr Leu Arg Lys Leu Ser Pro Asp Arg Phe Gln Arg Lys Arg Arg Leu Ser Gly Ala Gln Ala Pro Gly Pro Ser Val Pro Thr Arg Glu Pro Glu Gly Gly Thr Leu Ala Ala Trp Gln Glu Gly Glu Thr Glu Thr Ala Gln His Ser Gly

Leu Gly Leu Ile Asn Ser Phe Ala Ser Gly Glu Val Thr Thr Asn Gly Glu Ala Pro Glu Asn Gly Glu Asp Gly Glu His Gly Leu Leu Thr Tyr Ile Cys Asp Ala Met Glu Leu Gly Pro Gln Arg Ala Leu Lys Glu Glu Ser Gly Ala Lys Lys Lys Lys Asp Glu Glu Ser Lys Gln Gly Leu Arg Lys Pro Glu Leu Glu Lys Ala Ala Gln Ser Arg Arg Ser Ser Glu Asn Cys Ile Pro Ser Ser Asp Glu Pro Asp Ser Cys Gly Thr Gln Gly Pro Val Gly Val Glu Gln Val Gln Thr Gln Pro Arg Gly Arg Ala Ala Arg Gly Pro Gly Ser Ser Gly Thr Asp Ser Thr Arg Lys Pro Ala Ser Ala Val Gly Thr Pro Asp Lys Ala Gln Lys Ala Pro Gly Pro Gly Pro Gly Gln Glu Val Tyr Phe Ser Leu Lys Asp Met Tyr Leu Glu Asn Thr Gln Ala Val Arg Pro Leu Gly Glu Glu Gly Pro Gln Thr Leu Ser Val Arg Ala Pro Gly Glu Ser Pro Lys Gly Lys Ala Pro Leu Arg Ala Arg Ser Glu Gly Val Pro Gly Ala Pro Gly Gln Pro Thr His Ser Leu Thr Pro Gln Pro Thr Arg Pro Phe Asn Arg Lys Arg Phe Ala Pro Pro Lys Pro Lys Gly Glu Ala Thr Thr Asp Ser Lys Pro Ile Ser Ser Leu Ser Gln Ala Pro Glu Cys Gly Ala Gln Ser Leu Gly Lys Ala Pro Pro

Gln Ala Ser Val Gln Val Pro Thr Pro Pro Ala Arg Arg Arg His Gly 725 730 735

Thr Arg Asp Ser Thr Leu Gln Gly Gln Ala Gly His Arg Thr Pro Gly 740 745 750

Glu Val Leu Glu Cys Gln Thr Thr Thr Ala Pro Thr Met Ser Ala Ser 755 760 765

Ser Ser Ser Asp Val Ala Ser Ile Gly Val Ser Thr Ser Gly Ser Gln 770 780

Gly Ile Ile Glu Pro Met Asp Met Glu Thr Gln Glu Asp Gly Arg Thr 785 790 795 800

Ser Ala Asn Gln Arg Thr Gly Ser Lys Lys Asn Val Gln Ala Asp Gly 805 810 815

Lys Ile Gln Val Asp Gly Arg Thr Arg Gly Asp Gly Thr Gln Thr Ala 820 825 830

Gln Arg Thr Arg Ala Asp Arg Lys Thr Gln Val Asp Ala Gly Thr Gln 835 840 845

Glu Ser Lys Arg Pro Gln Ser Asp Arg Ser Ala Gln Lys Gly Met Met 850 860

Thr Gln Gly Arg Ala Glu Thr Gln Leu Glu Thr Thr Gln Ala Gly Glu 865 870 875 880

Lys Ile Gln Glu Asp Arg Lys Ala Gln Ala Asp Lys Gly Thr Gln Glu 885 890 895

Asp Arg Arg Met Gln Gly Glu Lys Gly Met Gln Gly Glu Lys Gly Thr 900 905 910

Gln Ser Glu Gly Ser Ala Pro Thr Ala Met Glu Gly Gln Ser Glu Gln 915 920 925

Glu Val Ala Thr Ser Leu Gly Pro Pro Ser Arg Thr Pro Lys Leu Pro 930 940

Pro Thr Ala Gly Pro Arg Ala Pro Leu Asn Ile Glu Cys Phe Val Gln 945 950 955 960

Thr Pro Glu Gly Ser Cys Phe Pro Lys Lys Pro Gly Cys Leu Pro Arg 965 970 975

Ser Glu Glu Ala Val Val Thr Ala Ser Arg Asn His Glu Gln Thr Val 980 985 990

Leu Gly Pro Leu Ser Gly Asn Leu Met Leu Pro Ala Gln Pro Pro His 995 1000 1005

Glu Gly Ser Val Glu Gln Val Gly Gly Glu Arg Cys Arg Gly Pro 1010 1015 1020

Gln Ser Ser Gly Pro Val Glu Ala Lys Gln Glu Asp Ser Pro Phe 1025 1030 1035

Gln Cys Pro Lys Glu Glu Arg Pro Gly Gly Val Pro Cys Met Asp. 1040 1050

Gln Gly Gly Cys Pro Leu Ala Gly Leu Ser Gln Glu Val Pro Thr 1055 1060 1065

Met Pro Ser Leu Pro Gly Thr Gly Leu Thr Ala Ser Pro Lys Ala 1070 1075 1080

Gly Pro Cys Ser Thr Pro Thr Ser Gln His Gly Ser Thr Ala Thr 1085 1090 1095

Phe Leu Pro Ser Glu Asp Gln Val Leu Met Ser Ser Ala Pro Thr 1100 1105 1110

Leu His Leu Gly Leu Gly Thr Pro Thr Gln Ser His Pro Pro Glu 1115 1120 1125

Thr Met Ala Thr Ser Ser Glu Gly Ala Cys Ala Gln Val Pro Asp 1130 1135 1140

Val Glu Gly Arg Thr Pro Gly Pro Arg Ser Cys Asp Pro Gly Leu 1145 1150 1155

Ile Asp Ser Leu Lys Asn Tyr Leu Leu Leu Leu Leu Lys Leu Ser 1160 1170

Ser Thr Glu Thr Ser Gly Ala Gly Gly Glu Ser Gln Val Gly Ala 1175 1180 1185

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Pro Arg Ala Glu Glu Leu Ala Ala Gly Asp Leu Gly Pro Ser Pro Lys Ala Gly Gly Leu Asp Thr Glu Val Ala Leu Asp Glu Gly Lys Gln Glu Thr Leu Ala Lys Pro Arg Lys Ala Lys Asp Leu Leu Lys Ala Pro Gln Val Ile Arg Lys Ile Arg Val Glu Gln Phe Pro Asp Ala Ser Gly Ser Leu Lys Leu Trp Cys Gln Phe Phe Asn Ile Leu Ser Asp Ser Val Leu Thr Trp Ala Lys Asp Gln Arg Pro Val Gly Glu Val Gly Arg Ser Ala Gly Asp Glu Gly Pro Ala Ala Leu Ala Ile Val Gln Ala Ser Pro Val Asp Cys Gly Val Tyr Arg Cys Thr Ile His Asn Glu His Gly Ser Ala Ser Thr Asp Phe Cys Leu Ser Pro Glu Val Leu Ser Gly Phe Ile Ser Arg Glu Glu Gly Glu 1575. Val Gly Glu Glu Ile Glu Met Thr Pro Met Val Phe Ala Lys Gly Leu Ala Asp Ser Gly Cys Trp Gly Asp Lys Leu Phe Gly Arg Leu Val Ser Glu Glu Leu Arg Gly Gly Gly Tyr Gly Cys Gly Leu Arg Lys Ala Ser Gln Ala Lys Val Ile Tyr Gly Leu Glu Pro Ile Phe Glu Ser Gly Arg Thr Cys Ile Ile Lys Val Ser Ser Leu Leu Val

Phe Gly Pro Ser Ser Glu Thr Ser Leu Val Gly Arg Asn Tyr Asp Val Thr Ile Gln Gly Cys Lys Ile Gln Asn Met Ser Arg Glu Tyr Cys Lys Ile Phe Ala Ala Glu Ala Arg Ala Ala Pro Gly Phe Gly Glu Val Pro Glu Ile Ile Pro Leu Tyr Leu Ile Tyr Arg Pro Ala Asn Asn Ile Pro Tyr Ala Thr Leu Glu Glu Asp Leu Gly Lys Pro Leu Glu Ser Tyr Cys Ser Arg Glu Trp Gly Cys Ala Glu Ala Pro Thr Ala Ser Gly Ser Ser Glu Ala Met Gln Lys Cys Gln Thr Phe Gln His Trp Leu Tyr Gln Trp Thr Asn Gly Ser Phe Leu Val Thr Asp Leu Ala Gly Val Asp Trp Lys Met Thr Asp Val Gln Ile Ala Thr Lys Leu Arg Gly Tyr Gln Gly Leu Lys Glu Ser Cys Phe Pro Ala Leu Leu Asp Arg Phe Ala Ser Ser His Gln Cys Asn Ala Tyr Cys Glu Leu Leu Gly Leu Thr Pro Leu Lys Gly Pro Glu Ala Ala His Pro Gln Ala Lys Ala Lys Gly Ser Lys Ser Pro Ser Ala Gly Arg Lys Gly Ser Gln Leu Ser Pro Gln Pro Gln Lys Lys Gly Leu Pro Ser Pro Gln Gly Thr Arg Lys Ser Ala Pro Ser Ser Lys Ala

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Lys Ser Glu Asp Gln Arg Cys Arg Ala Leu Leu Pro Ser Glu Leu Arg 35 40 45

Thr Leu Ile Gln Glu Ala Lys Glu Met Lys Trp Pro Phe Val Pro Glu 50 55 60

Lys Trp Gln Tyr Lys Gln Ala Val Gly Pro Glu Asp Lys Thr Asn Leu 70 75 80

Lys Asp. Val Ile Gly Ala Gly Leu Gln Gln Leu Leu Ala Ser Leu Arg 85 90 95

Ala Ser Ile Leu Ala Arg Asp Cys Ala Ala Ala Ala Ala Ile Val Phe 100 105 110

Leu Val Asp Arg Phe Leu Tyr Gly Leu Asp Val Ser Gly Lys Leu Leu 115 120 125

Gln Val Ala Lys Gly Leu His Lys Leu Gln Pro Ala Thr Pro Ile Ala 130 135 140

Pro Gln Val Val Ile Arg Gln Ala Arg Ile Ser Val Asn Ser Gly Lys 145 150 155 160

Leu Leu Lys Ala Glu Tyr Ile Leu Ser Ser Leu Ile Ser Asn Asn Gly 165 170 175

Ala Thr Gly Thr Trp Leu Tyr Arg Asn Glu Ser Asp Lys Val Leu Val 180 185 190

Gln Ser Val Cys Ile Gln Ile Arg Gly Gln Ile Leu Gln Lys Leu Gly 195 200 205

Met Trp Tyr Glu Ala Ala Glu Leu Ile Trp Ala Ser Ile Val Gly Tyr 210 215 220

Leu Ala Leu Pro Gln Pro Asp Lys Lys Gly Leu Ser Thr Ser Leu Gly 235 230 235

Ile Leu Ala Asp Ile Phe Val Ser Met Ser Lys Asn Asp Tyr Glu Lys 245 250 255

Phe Lys Asn Asn Pro Gln Ile Asn Leu Ser Leu Leu Lys Glu Phe Asp 260 265 270

His His Leu Leu Ser Ala Ala Glu Ala Cys Lys Leu Ala Ala Ala Phe 275 280 285

Ser Ala Tyr Thr Pro Leu Phe Val Leu Thr Ala Val Asn Ile Arg Gly 290 295 300

Thr Cys Leu Leu Ser Tyr Ser Ser Ser Asn Asp Cys Pro Pro Glu Leu 305 310 315 320

Lys Asn Leu His Leu Cys Glu Ala Lys Glu Ala Phe Glu Ile Gly Leu 325 330 335

Leu Thr Lys Arg Asp Asp Glu Pro Val Thr Gly Lys Gln Glu Leu His 340 345 350 $$

Ser Phe Val Lys Ala Ala Phe Gly Leu Thr Thr Val His Arg Arg Leu 355 360 365

His Gly Glu Thr Gly Thr Val His Ala Ala Ser Gln Leu Cys Lys Glu 370 375 380

Ala Met Gly Lys Leu Tyr Asn Phe Ser Thr Ser Ser Arg Ser Gln Asp 385 390 395 400

Arg Glu Ala Leu Ser Gln Glu Val Met Ser Val Ile Ala Gln Val Lys Glu His Leu Gln Val Gln Ser Phe Ser Asn Val Asp Asp Arg Ser Tyr Val Pro Glu Ser Phe Glu Cys Arg Leu Asp Lys Leu Ile Leu His Gly Gln Gly Asp Phe Gln Lys Ile Leu Asp Thr Tyr Ser Gln His His Thr Ser Val Cys Glu Val Phe Glu Ser Asp Cys Gly Asn Asn Lys Asn Glu Gln Lys Asp Ala Lys Thr Gly Val Cys Ile Thr Ala Leu Lys Thr Glu Ile Lys Asn Ile Asp Thr Val Ser Thr Thr Gln Glu Lys Pro His Cys Gln Arg Asp Thr Gly Ile Ser Ser Ser Leu Met Gly Lys Asn Val Gln Arg Glu Leu Arg Arg Gly Gly Arg Arg Asn Trp Thr His Ser Asp Ala Phe Arg Val Ser Leu Asp Gln Asp Val Glu Thr Glu Thr Glu Pro Ser Asp Tyr Ser Asn Gly Glu Gly Ala Val Phe Asn Lys Ser Leu Ser Gly Ser Gln Thr Ser Ser Ala Trp Ser Asn Leu Ser Gly Phe Ser Ser Ser Ala Ser Trp Glu Glu Val Asn Tyr His Val Asp Asp Arg Ser Ala Arg Lys Glu Pro Gly Lys Glu His Leu Val Asp Thr Gln Cys Ser Thr Ala Leu Ser Glu Glu Leu Glu Asn Asp Arg Glu Gly Arg Ala Met His Ser

Leu His Ser Gln Leu His Asp Leu Ser Leu Gln Glu Pro Asn Asn Asp

Asn Leu Glu Pro Ser Gln Asn Gln Pro Gln Gln Met Pro Leu Thr Pro Phe Ser Pro His Asn Thr Pro Gly Ile Phe Leu Ala Pro Gly Ala Gly Leu Leu Glu Gly Ala Pro Glu Gly Ile Gln Glu Val Arg Asn Met Gly Pro Arg Asn Thr Ser Ala His Ser Arg Pro Ser Tyr Arg Ser Ala Ser Trp Ser Ser Asp Ser Gly Arg Pro Lys Asn Met Gly Thr His Pro Ser Val Gln Lys Glu Glu Ala Phe Glu Ile Ile Val Glu Phe Pro Glu Thr Asn Cys Asp Val Lys Asp Arg Gln Gly Lys Glu Gln Gly Glu Glu Ile Ser Glu Arg Gly Ala Gly Pro Thr Phe Lys Ala Ser Pro Ser Trp Val Asp Pro Glu Gly Glu Thr Ala Glu Ser Thr Glu Asp Ala Pro Leu Asp Phe His Arg Val Leu His Asn Ser Leu Gly Asn Ile Ser Met Leu Pro Cys Ser Ser Phe Thr Pro Asn Trp Pro Val Gln Asn Pro Asp Ser Arg Lys Ser Gly Gly Pro Val Ala Glu Gln Gly Ile Asp Pro Asp Ala Ser Thr Val Asp Glu Glu Gly Gln Leu Leu Asp Ser Met Asp Val Pro Cys Thr Asn Gly His Gly Ser His Arg Leu Cys Ile Leu Arg Gln Pro Pro Gly Gln Arg Ala Glu Thr Pro Asn Ser Ser Val Ser Gly Asn Ile

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Gln Pro Gly Asn Met Leu Asn Cys Ser Gln Asn Ser Ser Ser Ser Ser 915 920 925

Val Trp Trp Leu Lys Ser Pro Ala Phe Ser Ser Gly Ser Ser Glu Gly 930 935 940

Asp Ser Pro Trp Ser Tyr Leu Asn Ser Ser Gly Ser Ser Trp Val Ser 945 950 955 960

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965 970 975

Pro Asp Asp Phe Glu Lys Leu Leu Ala Gly Val Arg His Asp Trp Leu 980 985 990

Phe Gln Arg Leu Glu Asn Thr Gly Val Phe Lys Pro Ser Gln Leu His 995 1000 1005

Arg Ala His Ser Ala Leu Leu Leu Lys Tyr Ser Lys Lys Ser Glu 1010 1015 1020

Leu Trp Thr Ala Gln Glu Thr Ile Val Tyr Leu Gly Asp Tyr Leu 1025 1030 1035

Thr Val Lys Lys Gly Arg Gln Arg Asn Ala Phe Trp Val His 1040 1045 1050

His Leu His Gln Glu Glu Ile Leu Gly Arg Tyr Val Gly Lys Asp 1055 1060 1065

Tyr Lys Glu Gln Lys Gly Leu Trp His His Phe Thr Asp Val Glu 1070 1080

Arg Gln Met Thr Ala Gln His Tyr Val Thr Glu Phe Asn Lys Arg 1085 1090 1095

Leu Tyr Glu Gln Asn Ile Pro Thr Gln Ile Phe Tyr Ile Pro Ser 1100 1105 1110

Thr Ile Leu Leu Ile Leu Glu Asp Lys Thr Ile Lys Gly Cys Ile 1115 1120 1125

Ser Val Glu Pro Tyr Ile Leu Gly Glu Phe Val Lys Leu Ser Asn 1130 1135 1140 Asn Thr Lys Val Val Lys Thr Glu Tyr Lys Ala Thr Glu Tyr Gly 1145 1150 1155 Leu Ala Tyr Gly His Phe Ser Tyr Glu Phe Ser Asn His Arg Asp 1160 1170 1165 Val Val Asp Leu Gln Gly Trp Val Thr Gly Asn Gly Lys Gly 1175 1180 1185 Leu Ile Tyr Leu Thr Asp Pro Gln Ile His Ser Val Asp Gln Lys 1190 1195 1200 Val Phe Thr Thr Asn Phe Gly Lys Arg Gly Ile Phe Tyr Phe Phe 1205 1210 1215 Asn Asn Gln His Val Glu Cys Asn Glu Ile Cys His Arg Leu Ser 1220 1225 1230 Leu Thr Arg Pro Ser Met Glu Lys Pro 1235 1240 <210> 42 <211> 30 <212> DNA<213> Artificial Sequence <220> <223> primer <400> 42 30 ctgcgacaga gactacatgg ggtagaactc <210> 43 <211> 30 <212> DNA <213> Artificial Sequence <220> <223> primer <400> 43 tgagtgtctt cggtagatgg ccttctactg 30 <210> 44 <211> 21 <212> DNA <213> Artificial Sequence <220> <223> primer

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